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doi:10.1074/jbc.271.42.26209

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	&isopropylmalatereplicates autonomously, whereas strain AH22/CV9-2 carries multiple copies of theabout 50 copies/cell in many strains of S. cerevisiae (IO, 17). Strain AH22/CV9-2 was obtained from T. D. Petes					
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	•1980), and the yeast <b>S</b> . cerevisiae. These included the	erevis	iae (Storms et al., 1984intervening se nally to all spliced <b>S. cerevisiae</b> genes a AH22 MATa leu2-3,112 his4-419		View Shopping Cart	

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Molecular Engineering Study on Electron Transfer from NADPH-P450

Reductase to Rat Mitochondrial P450c27 in Yeast Microsomes

J. Biol. Chem., Oct 1996; 271: 26209 - 26213;

Saccharomyces cerevisiae AH22 (9) were usedtransformation of E.coli and S.cerevisiae cells were performedRecombinant E.coli and S.cerevisiae strains were cultivatedwas purified from AH22/pAMS25 microsomesintroduced into S.cerevisiae AH22 cells to obtain	
NUCLEIC ACIDS, PROTEIN SYNTHESIS, AND MOLECULAR GENETICS: Yoon Lee and Ross N. Nazar Ribosomal 5 S rRNA Maturation in Saccharomyces cerevisiae  J. Biol. Chem., Jun 1997; 272: 15206 - 15212; doi:10.1074/jbc.272.24.15206 transform a LEU2-deficient yeast strain (AH22) as described by Hinnen et alBlobel () with minor modifications. S. cerevisiae, strain AH22, was grown at 30C in YEPD medium (1sequences on the expression of 5 S rRNA in S. cerevisiae. The 5 S rRNA genes containing altered	Abstract Full Text PDF Purchase Article View Shopping Cart
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DI Van Ryk, Y Lee, and RN Nazar  Unbalanced ribosome assembly in Saccharomyces cerevisiae  expressing mutant 5 S rRNAs  J. Biol. Chem., Aug 1992; 267: 16177 - 16181 transform a Leu-2-deficient yeast strain (AH22) by the method of Hinnen et al.  (Hinnen16178 Unbalanced Ribosome Assembly in S. cerevisiae orthophosphate were added, and at appropriateconditions Unbalanced Ribosome Assembly in S. cerevisiae 16179 z 0 i;:kk 2 0.5 v)  O 1 10 20Polyribosome profiles in transformed S. cerevisiae expressing mutant 5 S rRNA genes.  Polyribosomes	Abstract PDF Purchase Article View Shopping Cart
DI Van Ryk, Y Lee, and RN Nazar  Efficient expression and utilization of mutant 5 S rRNA in  Saccharomyces cerevisiae  J. Biol. Chem., May 1990; 265: 8377 - 8381	Abstract PDF Purchase Article View Shopping Cart
Mutants of Saccharomyces cerevisiae defective in sn-glycerol-3-phosphate acyltransferase. Simultaneous loss of dihydroxyacetone phosphate acyltransferase indicates a common gene  J. Biol. Chem., Jul 1986; 261: 9144 - 9149 acyltransferase activity of s. cerevisiae. This paper reports theSigma. Palmi9144 Mutants in S. cerevisiae sn-Glycerol-3-phosphateGenetic Stock Center. Strains AH22 (a, leu2-3, leu2-112, his4-519could be due to Mutants in S. cerevisiae sn-Glycerol-3-phosphateTTAI was mated with strain AH22, the diploids sporulatedacyltransferase. 9 148 Mutants in S. cerevisiae sn-Glycerol-3-phosphate	Abstract PDF Purchase Article View Shopping Cart
NUCLEIC ACIDS, PROTEIN SYNTHESIS, AND MOLECULAR GENETICS: Yoon Lee, Yuri F. Melekhovets, and Ross N. Nazar Termination as a Factor in `Quality Control'' during Ribosome Biogenesis J. Biol. Chem., Nov 1995; 270: 28003 - 28005; doi:10.1074/jbc.270.47.28003 Saccharomyces cerevisiae strain (AH22) as described by Hinnen etAltered 3`-ETS regions in the S. cerevisiae 5 S rRNA gene (Fig. 2) wereAltered 3 9-ETS regions in the S. cerevisiae 5 S rRNA gene (Fig. 2) weretransform LEU2-deficient S. cerevisiae cells. Whole cell RNA wasSaccharomyces cerevisiae strain (AH22) as described by Hinnen et	Abstract Full Text PDF Purchase Article View Shopping Cart
WS Moye, N Amuro, JK Rao, and H Zalkin  Nucleotide sequence of yeast GDH1 encoding nicotinamide adenine dinucleotide phosphate-dependent glutamate dehydrogenase  J. Biol. Chem., Jul 1985; 260: 8502 - 8508 training and sequenced the S. cerevisiae GDHI gene encoding  NADPdependentcomplementation in bacteria. Yeast strain AH22 was transformed with pGDH1this transformant relative to AH22 transformed with the cloningThe E. coli, N. crassa, and S. cerevisiae enzymes have 189 conserved	• Abstract • PDF • Purchase • Article • View • Shopping Cart

	Mutants of Saccharomyces cerevisiae defective in sn-1,2-diacylglycerol cholinephosphotransferase. Isolation, characterization, and cloning of the CPT1 gene  1. Biol. Chem. Mar 1987: 262: 3909 - 3917	► Abstract ► PDF Purchase Article View ► Shopping Cart
	Unbalanced regulation of the ribosomal 5 S RNA-binding protein in Saccharomyces cerevisiae expressing mutant 5 S rRNAs  J. Biol. Chem., Sep 1992; 267: 17738 - 17742	> Abstract > PDF Purchase Article View > Shopping Cart
	T Kodaki and S Yamashita  Yeast phosphatidylethanolamine methylation pathway. Cloning and characterization of two distinct methyltransferase genes  J. Biol. Chem., Nov 1987; 262: 15428 - 15435  ▶1.1.17 CHO2 protein, S cerevisiae   EC 2.1.1.17 PhosphatidylethanolamineUniversity of California). AH22 (a leu23 leu2-112 his4-519crossing D159-6B with AH22. Strain 301 (a pemlcrossing strain 314 with AH22. Bacterial Strains andidentified two types of S. cerevisiae mutants defective in	➤ Abstract ➤ PDF Purchase Article View ➤ Shopping Cart
	B Seraphin, M Simon, and G Faye  The mitochondrial reading frame RF3 is a functional gene in  Saccharomyces uvarum  J. Biol. Chem., Jul 1987; 262: 10146 - 10153 X4004-3A FF314 FLIOO M12-54 X2180-1A KL14-4A AH22 GRF18 AB320 S288C MH41-7B GRF18/2 BS104-1codon, respectively. This confirms that S. cerevisiae and S. uvarum are phylogenetically relatedcytoduction (Lancashire and Mattoon, 1979) to S. cerevisiae and have no effect on growth and respiration	Abstract PDF Purchase Article View Shopping Cart
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	MP Nobrega, FG Nobrega, and A Tzagoloff COX10 codes for a protein homologous to the ORF1 product of Paracoccus denitrificans and is required for the synthesis of yeast cytochrome oxidase  J. Biol. Chem., Aug 1990; 265: 14220 − 14226  →EXPERIMENTAL PROCEDURES Strains and Media-The strains of S. cerevisiae used in this study are listed in Table I. The ~0x10 mutantsW303-1B a,p+,ade2-1,his3-11,3-15,ku2-3,2-112,trp1- 1,ura3-1 AH22 a,p+,bu2-3,2-112,his4-519 C262 a,p+,met6,coxlO-4,ripl C26219 R. Rothstein" R. Rothstein" G. Fink" This study C262 x AH22 This study C22 x CBll B22 x LH20 Ref. 17 This study This study	► Abstract ► PDF ► Purchase Article View ► Shopping Cart
	M Iwasaki, R Juvonen, R Lindberg, and M Negishi  Alteration of high and low spin equilibrium by a single mutation of amino acid 209 in mouse cytochromes P450  J. Biol. Chem., Feb 1991; 266: 3380 - 3382 these mutated cDNAs was inserted into the yeast expression vector pAAH5 as described previously (6) and then transformed to S. cerevisiae AH22 cells using the LiCl method. Partially, Purification of Mutated P450s-Microsomes were prepared from a 10-liter culture	► Abstract ► PDF Purchase Article View ► Shopping Cart
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